

The Paradigm of Huntington's Disease: Therapeutic Opportunities in Neurodegeneration

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Summary: Despite a relatively small number of affected patients, Huntington's disease (HD) has been a historically important disease, embodying many of the major themes in modern neuroscience, including molecular genetics, selective neuronal vulnerability, excitotoxicity, mitochondrial dysfunction, apoptosis, and transcriptional dysregulation. The discovery of the *HD* gene in 1993 opened the door to the mechanisms of HD pathogenesis.

Multiple pathologic mechanisms have been discovered, each one serving as a potential therapeutic target. HD thus continues to serve as a paradigmatic disorder, with basic bench research generating clinically relevant insights and stimulating the development of therapeutic human trials. **Key Words:** Huntington's disease, apoptosis, excitotoxicity, mitochondria, transcription, transplantation.

INTRODUCTION

In many ways, Huntington's disease (HD) has been a model disorder. HD was the first autosomal dominant disorder in which the then-novel technique of "reverse genetics" was successfully applied.¹⁻⁴ It would be another 10 years until the gene was found. Excitotoxicity, the idea that overexcitation could lead to the death of neurons, was first seriously considered in the context of HD.⁵⁻⁸ Excitotoxicity has now been implicated in a host of neurologic disorders.⁹⁻¹¹ Similarly, many of the major ideas in modern neuroscience, including mitochondrial dysfunction, apoptosis, and transcriptional dysregulation, were initially derived from the HD field or have been explored substantially in HD. When the *HD* gene was eventually identified as a CAG trinucleotide repeat disorder, HD joined a novel class of neurodegenerative diseases, the polyglutamine disorders.^{12,13} The discovery of intracellular aggregates of mutant huntingtin bolstered the emerging concept that all neurodegenerative disorders are diseases of protein misfolding.¹⁴⁻¹⁷

Clinically, HD has also been a paradigmatic disease. As for so many neurologic diseases, there is no effective therapy for HD, which remains a progressive, fatal dis-

order. However, just as advances in molecular genetics made possible predictive genetic testing for HD, recent laboratory discoveries are pointing the way to novel therapeutic approaches.¹⁸ Large multi-center clinical HD trials have been completed, and many more trials are planned. In this review, we outline recent advances in understanding the molecular pathogenesis of HD, as well as the therapeutic opportunities created by such advances.

OVERVIEW OF HUNTINGTON'S DISEASE

Clinical features

HD is a neurodegenerative disease characterized by the clinical triad of movement disorder, dementia and psychiatric disturbance.¹⁹ With an incidence of 4 to 10 per 100,000, HD afflicts 30,000 people in the United States. Another 250,000 persons are genetically at risk. HD is inherited in an autosomal dominant manner and typically develops in the fourth or fifth decade of life. Initially, patients demonstrate personality changes and develop small involuntary movements. As the disease progresses, the movement disorder becomes more pronounced, and cognitive deficits as well as psychiatric disturbances occur. Though the movement disorder is typically chorea, virtually any type of movement disorder is seen, including dystonia, rigidity, myoclonus, and athetosis. Cognitive dysfunction can be due to striatal or cortical degeneration and includes dementia and difficulties with executive functioning. Psychiatric disturbances

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most commonly manifest as apathy and depression but obsessive-compulsive disorder, psychosis, paranoia, and substance abuse also occur.²⁰

Disease duration is 10 to 30 years, and later symptoms include rigidity, dystonia, and bradykinesia. Weight loss is a common feature of the disease. Death usually results from aspiration pneumonia secondary to dysphagia, or from complications resulting from falls or chronic illness; suicide is another cause of death. Currently, there is no effective treatment to delay the onset or slow the progression of HD.

A juvenile form of HD exists and is most often inherited through paternal transmission. The movement disorder tends to be more parkinsonian than the adult form and is characterized by bradykinesia, rigidity, and resting tremor. Seizures are also common. Patients with juvenile onset of the disease have a more severe course of disease with an average duration of 5 to 20 years.

Pathology

Pathologically, HD is characterized by the selective loss of efferent medium spiny neurons in the striatum (caudate nucleus and putamen) of the basal ganglia.²¹ The striatum is comprised of numerous neuronal subtypes which include medium spiny projection neurons and interneurons. The latter category encompasses the medium spiny reduced nicotinamide adenine dinucleotide phosphate (NADPH)-diaphorase-positive, neuronal nitric oxide synthase (nNOS)-positive neurons and large spiny cholinergic neurons. Unlike the medium spiny projection neurons, interneuron populations are relatively spared in HD.^{22–26} The factors that render striatal projection neurons more susceptible to damage are unknown.

Although the striatum is the most profoundly affected region in HD, degenerative changes in the cortex, thalamus, and subthalamic nucleus have also been reported in HD.^{27,28} Therefore, the changes in the striatum reflect a relative selective vulnerability to cell death. One possible explanation is that other brain regions in HD are affected, but at a slower rate than the striatum. In juvenile HD, in which the rate of disease progression is much faster, cortical pathology is more significant.

The pathologic changes in HD appear to precede the frank appearance of disease symptoms. Decreases in mRNA encoding dopamine D1 and D2 receptors have been found in the striatum of patients in the early stages of HD.²⁹ Positron emission tomography studies of gene-positive but clinically asymptomatic patients demonstrate that dopamine D1 and D2 receptors are decreased before the onset of symptoms.^{30–32} In transgenic mouse models of HD, the alterations in brain receptor expression precede the development of obvious neurological symptoms.^{33–35} Thus, neuronal dysfunction predates the appearance of neurologic symptomatology in HD.

HD is a polyglutamine disorder

Although a chromosomal marker was identified in 1983,³ it was not until 1993 that the *HD* gene was discovered.³⁶ The *IT15* ("interesting transcript 15") gene encodes a 348-kDa protein called huntingtin whose amino acid sequence does not resemble that of any known protein. The 5' end of the gene contains a polymorphic trinucleotide repeat (cytosine-adenine-guanosine, CAG) which encodes a series of glutamines. Normal persons have CAG repeat lengths of 7–34. The CAG repeat is expanded and unstable in HD patients, with repeat length inversely correlating with age of disease onset. Repeat lengths of >40 glutamines invariably produce HD, and repeats of >100 glutamines invariably cause juvenile onset.³⁷ CAG repeat lengths of 35–39 are considered intermediate alleles.³⁸

By virtue of having an expanded polyglutamine moiety, HD becomes a member of a novel class of neurodegenerative disorders known as CAG trinucleotide repeat disorders or polyglutamine disorders.^{12,39,40} Although some trinucleotide disorders such as myotonic dystrophy or fragile X mental retardation have expansions of repeats in noncoding regions of the gene, the CAG repeat disorders are distinct. In these disorders, the expansion of the CAG repeat (the codon encoding the amino acid glutamine) occurs within the coding region of the gene, and the disease gene is translated into protein. The polyglutamine disorders include spinobulbar muscular atrophy (SBMA or Kennedy's disease), dentatorubropallidoluysian atrophy (DRPLA), and several of the spinocerebellar ataxias (SCA-1, -2, -3, -6, -7, -12, and -17). Although these are relatively rare disorders, they share some remarkable features. All are autosomal dominant or sex-linked dominant diseases, usually with adult onset, characterized by progressive neurodegeneration of selected neuronal populations. For all polyglutamine disorders, higher numbers of repeats in the mutant gene correlates with an earlier age of onset. Polyglutamine protein-containing inclusions are another striking common feature, which are detected in human pathologic specimens as well as in transgenic mouse models. Interestingly, although all of the mutant proteins share the polyglutamine region, the other portions of the host proteins are not homologous. Given the remarkable similarities between these disorders, these observations point to the polyglutamine moiety as the most relevant portion of the molecular contributing to disease pathogenesis.⁴¹

Two experiments involving transgenic mice confirm the primary importance of the polyglutamine tract in disease pathogenesis. Bates and colleagues⁴² created lines of mice with a transgene containing only exon 1 of the human *HD* gene. While exon 1 contains the polymorphic CAG region, there are 67 exons in total, and thus exon 1 comprises only about 3% of the complete gene. Remarkably, transgenic mice expressing a mutated

exon 1 develop an abnormal neurologic phenotype that is reminiscent of HD: normal initial development, but later onset of motor and cognitive difficulties, weight loss, and early death.⁴² In another experiment, Ordway and colleagues⁴³ inserted CAG repeats into the mouse *HPRT* gene, which normally does not contain CAG repeats. Transgenic mice expressing these “ectopic” repeats developed an abnormal motor phenotype and ubiquitinated intranuclear inclusions, features which have been observed in a numbers of transgenic mouse models of polyglutamine diseases. While these observations certainly do not exclude a role for the remainder of the huntingtin molecule, they highlight the primary pathologic relevance of the expanded polyglutamine portion of the protein.

Function of huntingtin

Huntingtin itself has no clear homology to known proteins, and its normal function is not known. Huntingtin is certainly required for normal development, as knockout mice lacking their endogenous huntingtin die at embryonic stage.^{44,45} Mutant forms of huntingtin can rescue the embryonic lethality of the knockout mouse, indicating that the polyglutamine expansion does not remove all of the normal function of huntingtin.⁴⁶ This observation also bolsters the idea that HD is a gain of function disease. Huntingtin is also required for normal hematopoiesis and spermatogenesis.^{46,47}

Recent data support an anti-apoptotic role for normal (wild-type) huntingtin.^{48,49} Cattaneo and colleagues⁵⁰ have found that wild-type huntingtin exerts its anti-apoptotic effects by preventing the processing of caspase-9. Wild-type huntingtin also promotes transcription of brain-derived neurotrophic factor (BDNF).⁵¹ Since wild-type huntingtin is itself cleaved in HD tissues,^{51,52} some of the features of HD may be the result of loss of function of wild-type huntingtin. Caspases have been implicated in the cleavage of both mutant and wild-type huntingtin.^{53,54} Wild-type huntingtin can also be cleaved by calpain,^{55,56} and an aspartyl protease.⁵⁷ While proteolytic cleavage of wild-type huntingtin could lead to a loss of normal function, cleavage into smaller fragments also alters the nuclear trafficking of huntingtin.^{58–61} While huntingtin is predominantly cytoplasmic, a small fraction of full-length huntingtin is also normally found in the nucleus.^{62,63}

THEORIES OF PATHOGENESIS

The exact mechanism by which neurons die in HD remains unknown. The discovery of the *HD* gene, and, more recently, the development of transgenic mouse, fly, and worm models has led to an explosion of discovery of the pathogenic mechanisms caused by mutant huntingtin. Far from arriving at a unifying mechanism, there is in-

stead robust evidence that multiple pathologic mechanisms occur in HD.

Excitotoxicity

The death of neurons as a result of overactivity of glutamate neurotransmission, a process termed excitotoxicity, has been an influential theme in modern neuroscience. Much of the support for the excitotoxic hypothesis has come from the HD field.⁶⁴ The theory of excitotoxicity as a pathogenic mechanism in HD has emerged over the last few decades, beginning with the observation that injections of excitatory amino acids into the striatum of rodents and primates led to neuronal depletion and a neurologic phenotype that was similar to HD. As the striatum receives large glutamatergic input from corticostriatal afferents, it is a structure at risk of glutamate-mediated excitotoxic injury. Intrastriatal injections of glutamate agonists, particularly those acting at NMDA receptors, have been used to create animal models of HD. Of the NMDA agonists, quinolinic acid, an endogenous metabolite of tryptophan, has been shown to most closely mimic the neuropathology and neurologic phenotype of HD, producing selective degeneration of medium spiny neurons with sparing of interneuron populations. In addition, glutamate receptors have been shown to be reduced in human HD brains.^{67–69} In more recent studies, subsets of glutamate receptors, specifically the metabotropic glutamate receptor mGluR2, appear to be significantly decreased in a transgenic mouse model of HD.³⁴ mGluR2 receptors down-regulate glutamate release at corticostriatal presynaptic terminals, and depletion of these receptors could lead to overstimulation of postsynaptic striatal neurons. Further data supporting the excitotoxic model show that there is increased sensitivity to NMDA receptor activation and enhanced excitotoxicity in HD transgenic mice.^{70–73} Finally, perturbations in glutamate handling in the brain may also contribute to excitotoxicity in HD as there appears to be down-regulation of the GLT-1 glial glutamate transporter in a transgenic mouse model of HD.⁷⁴

In addition, mutant huntingtin may alter the structure of the postsynaptic apparatus. Recent data demonstrate an association in human cortex between huntingtin and PSD-95 (post-synaptic density protein 95, a protein that is involved in the post-synaptic clustering of NMDA receptors) that is mediated by the SH3 domain of PSD-95. The presence of the polyglutamine expansion in mutant huntingtin appears to significantly reduce binding to PSD-95 and promotes sensitization of NMDA receptors. These data raise the intriguing possibility that wild-type huntingtin may function to modulate NMDA receptor activity by dissociating NMDA receptor-linked macromolecular complexes.⁷⁵

Although the excitotoxic hypothesis is a persuasive one, there are still some features of HD that remain

unexplained. For example, in addition to the striatum, the hippocampus, cortex, and cerebellum contain similar or higher levels of glutamate receptors, yet there is a strikingly selective loss of striatal neurons in HD. Thus, the high-level glutamate receptors cannot explain the regional specificity of HD pathology. One possibility is that the selective loss of striatal neurons is due to differences in glutamate receptor subtypes in these brain regions. The cell selectivity within the striatum remains similarly unexplained. Recent data suggest that regional expression of the NR2B receptor subunit of the NMDA receptor accounts for the severity of neuronal death in HD.^{76,77} Mutant huntingtin also selectively increases current flow through NMDA receptors comprised of NR1/NR2B subunits.^{72,73,78}

Apoptosis/caspases

One of the major ideas in modern neuroscience is that neurons may die by inappropriately activating the apoptosis cell death program.⁷⁹ Within the last decade, the role of apoptosis and caspases, the apoptosis-related cysteine proteases, has been proposed in the pathogenesis of HD.^{52,80,81} Apoptosis, also known as programmed cell death, is a feature of a number of chronic neurodegenerative diseases [Alzheimer's disease, amyotrophic lateral sclerosis (ALS), HIV dementia], and caspases are important in initiating and executing the cell death program. Recent data have shown that there is increased activation of caspase-1 in pre-symptomatic and early symptomatic HD transgenic mice.⁵² In cellular model systems, expression of mutant huntingtin can induce apoptosis.^{61,82,83} The normal and mutant huntingtin proteins are substrates for caspase-1 and caspase-3, and both are cleaved, generating N-terminal fragments.^{53,84,85} The mutant huntingtin N-terminal fragments, which contain the expanded polyglutamine moiety, are translocated to the nucleus, where they accumulate into aggregates. The presence of these mutant N-terminal fragments in the nucleus is a stimulus for caspase-1 up-regulation.⁸³ As the disease progresses, caspase-3, -8, and -9 are activated, and there is release of cytochrome *c*, which serves as an apoptotic trigger.⁸⁶ Inhibition of caspase-1 appears to slow disease progression in HD transgenic mice; transgenic HD mice also expressing a dominant-negative caspase-1 gene survived longer and had delayed appearance of symptoms, neuronal inclusions, and neurotransmitter receptor changes.⁵² The activation of caspases has also been shown in human HD striatal tissue.^{52,86,87}

Mitochondrial dysfunction

Neurons are metabolically active cells; processes that affect mitochondrial function have disproportionate effects on neurons. Mitochondrial dysfunction has been proposed to underlie a number of neurodegenerative disorders, including HD.⁸⁸⁻⁹¹ Defects in energy metabolism have been well-documented in the disorder: glucose me-

tabolism is decreased in the brains of HD patients,⁹² lactate levels are increased in areas of HD brain,⁹³ and the cerebrospinal fluid lactate/pyruvate ratio in HD patients is increased as compared with age-matched controls.⁹⁴ In addition, metabolic defects in skeletal muscle have been reported,^{94,95} a finding consistent with observations that HD patients suffer progressive weight loss despite increased caloric intake. Thus, defects in energy metabolism appear to be widespread in HD, affecting both the brain and peripheral tissues.

Reports of changes in activity of enzymes involved in oxidative phosphorylation lend further support to the mitochondrial dysfunction hypothesis in HD. Reduced activity of the complex II enzyme succinate dehydrogenase (SDH) has been found in postmortem HD brain tissue.⁹⁶ In addition, a decrease in complex II-III activity in HD basal ganglia has been observed.^{97,98} Systemic administration of the mitochondrial toxin 3-nitropropionic acid (3-NP), an irreversible inhibitor of SDH, to nonhuman primates produces a movement disorder and cognitive deficits similar to HD. Histologic examination of the striatum 3-NP-treated animals reveals a pattern of neuronal loss similar to HD with abnormalities in medium spiny neurons and sparing of the NADPH diaphorase interneurons and nucleus accumbens.^{99,100} Malonate, another complex II inhibitor, has also been used to produce such "chemical lesion" models of HD.¹⁰¹

The mechanism by which mitochondrial dysfunction ultimately leads to neuronal death has been proposed to be linked to excitotoxicity because NMDA receptor antagonists can block the effect of mitochondrial inhibitors.¹⁰² Recent data show that 3-NP treatment induces long-term NMDA-mediated excitation in medium spiny neurons.¹⁰³ These observations lend credence to the "weak excitotoxic hypothesis" in which mitochondrial inhibition functions to tip the excitotoxic balance toward neurotoxicity.¹⁰⁴

Recently, Panov and colleagues¹⁰⁵ have demonstrated that mitochondria isolated from HD patient lymphoblasts have reduced calcium buffering capacity. In addition to being able to take up less ionized calcium than wild-type mitochondria, HD mitochondria are relatively depolarized. Since the mitochondrial membrane potential is directly related to ATP production, reduced membrane potential likely contributes to energetic failure in this disease. Interestingly, in an HD knock-in mouse model, there is a progressive reduction in the amount of cyclic AMP, indicating an energy deficit.¹⁰⁶

While derangements in energy metabolism are well-documented in HD, it is still unclear whether these defects are a cause of the disease or a consequence. Some studies have shown that disorders in metabolism are found in both presymptomatic and symptomatic HD individuals (glucose, lactate, and muscle). However, other studies have shown no evidence of perturbations in mi-

tochondrial transport in the brains of postmortem HD patients and in transgenic mice expressing full-length mutant huntingtin.¹⁰⁷ Therefore the exact role of mitochondrial dysfunction in the pathogenesis of HD has yet to be determined.

Transcriptional dysregulation

The identification of the *HD* gene has led to new theories regarding pathogenesis. As detailed above, huntingtin contains a polyglutamine stretch, the expansion of which confers a gain-of-function property. Polyglutamine tracts are a feature of transcription factor proteins (e.g., Sp1, TBP, TRAM1, and CBP) and can act as activation domains for transcription factors *in vitro*. Mutant huntingtin has been shown to have an aberrant nuclear localization as well as altered protein-protein interactions, and the combination of these observations suggests mutant huntingtin might act to sequester polyglutamine-containing nuclear transcription factors, thereby leading to transcriptional dysregulation and subsequent neuronal death.¹⁰⁸ Recent data have shown reduced expression of a number of genes in a transgenic mouse model of HD^{33,34,109,110} suggesting that transcriptional repression is the main result of dysregulation. In addition, a number of studies have demonstrated that huntingtin is able to bind transcription factors resulting in decreased activity of these proteins.^{111–113} Not surprisingly, many of the transcription factors with altered activity in HD directly or indirectly regulate histone acetylation, a process which helps regulate transcription through covalent modification of chromatin.

NEUROTHERAPEUTICS IN HD

Currently there is no cure for HD, and there are no therapies which significantly slow the progression of the disease or delay its onset. However, basic science research into the pathogenesis of HD has led to a number of different and promising areas of neurotherapeutics. One significant advantage in the study of HD is the proliferation of animal and invertebrate disease models which have aided not only in investigating the function of huntingtin, but also in screening and testing potential therapeutic drugs.

Excitotoxicity

The excitotoxic hypothesis is a well-studied mechanism in HD pathology and has led to the identification of possible neuroprotective compounds. As detailed above, injection of quinolinic acid into the striatum of rodents recapitulates many of the features of HD and has therefore been used as a model to identify and test potential neuroprotective agents. N-acetyl-aspartyl-glutamate, a compound which has antagonist (and agonist) activity at NMDA receptors and agonist activity at the mGluR3 metabotropic glutamate receptor, has been found to re-

duce lesion volume in quinolinic-acid-treated rats.¹¹⁷ (S)-4-Carboxy-3-hydroxyphenylglycine, which acts at metabotropic glutamate receptors, has also been shown to protect against excitotoxicity in rats that have received intrastriatal injection of quinolinic acid.¹¹⁸

Riluzole, an inhibitor of glutamate release that is used for ALS, was found to be protective against quinolinate lesions.¹¹⁹ An open-label human trial found that riluzole reduced chorea and showed decreased levels of cerebral lactate, as assessed by magnetic resonance spectroscopy.¹²⁰ Another open-label trial of riluzole also demonstrated transient motor improvement in human HD subjects.¹²¹ Riluzole has also been studied in a transgenic mouse model of HD.¹²² Mice who received riluzole had increased lifetime with delay in weight loss. Neuropathologically, the striata of riluzole-treated mice revealed fewer and smaller nuclear aggregates than the striata of untreated transgenic mice. Interestingly, riluzole has also been found to up-regulate the levels of several key neurotrophic factors, including BDNF and glial-derived neurotrophic factor (GDNF).¹²³ A large multi-center placebo-controlled human clinical trial of riluzole is currently underway in Europe.

The recent finding that specific NMDA receptor subtypes (those comprised of NR1A/NR2B subunits) are partly responsible for the selective striatal neuronal vulnerability in HD suggests that subunit-specific antagonists may have a therapeutic role in HD. Ifenprodil, an NR2B-specific antagonist, has been shown to eliminate excitotoxic cell death in medium spiny neurons from transgenic and wild-type mice after exposure to NMDA.⁷³

A recent multi-center trial called (CARE-HD) (Coenzyme Q10 and Remacemide in Huntington's Disease) tested the effects of coenzyme Q10, an antioxidant and cofactor involved in mitochondrial electron transfer, and remacemide, a noncompetitive NMDA receptor antagonist.¹²⁴ In the largest randomized, placebo-controlled clinical HD trial to date, 348 patients were treated for 24 months with either coenzyme Q10, remacemide, a combination of both drugs, or none. In two different lines of transgenic mice, the combination of these two compounds demonstrated benefit.^{125,126} However, in the CARE-HD trial, there was no evidence of any benefit of remacemide, either alone or in combination with coenzyme Q10.

In addition to glutamate receptor subtypes, other receptor subtypes have been implicated in the pathogenesis of HD. Defects in the locus coeruleus-noradrenergic system have been observed in HD and the blockade of the presynaptic inhibitory α 2-adrenergic receptors has been studied in the quinolinate lesion model. Administration of efaroxan and idazoxan, two α 2-adrenergic receptor antagonists, to quinolinate-injected mice was found to decrease certain behavioral and biochemical characteris-

tics of HD, specifically ipsiversive circling responses to apomorphine and choline acetyltransferase deficit in the affected striatum.¹²⁷

Dopamine has been postulated to be involved in neurotoxicity seen in HD.¹²⁸ Transgenic HD mice have reduced levels of dopamine¹²⁹; numerous studies have found decreased dopamine receptors in both human and transgenic mouse models of HD.¹³⁰ Transgenic HD mice given L-dopa experience transient improvement in motor symptoms, but later show signs of increased toxicity.¹³¹

Mitochondrial toxicity

Molecules that function to increase energy by boosting ATP stores have been studied for their potential neuroprotective effects. Both creatine and cyclocreatine have been investigated in animal models of HD. In rats treated with 3-NP, oral creatine supplementation produced significant protection against malonate- and 3-NP-induced lesions.¹³² Cyclocreatine protected against malonate-induced but not 3-NP lesions. The neuroprotective effects of creatine have also been demonstrated in a transgenic HD mice.¹³³ Transgenic R6/2 mice given oral supplementation with creatine exhibited improved survival, delayed atrophy of striatal neurons and delayed formation of huntingtin-positive aggregates. In addition, body weight was significantly greater in creatine-treated transgenic mice and motoric ability, as measured by performance of the Rotarod test, was significantly improved in those animals. These data support a role for metabolic dysfunction as a pathogenic component of HD and suggest a potential role for ATP repletion via creatine supplementation as a therapeutic strategy. Human clinical trials of creatine in HD patients are underway.

Other studies have investigated the therapeutic potential of antioxidants. Both vitamin E and idebenone have been tested in clinical trials and found to have no significant impact on functional decline in HD.^{134,135} OPC-14117, a lipophilic free-radical scavenger that concentrates in the brain has been tested in a safety and tolerability trial.¹³⁶ The CARE-HD trial tested the effects of coenzyme Q10, an antioxidant and cofactor involved in mitochondrial electron transfer, as well as remacemide.¹²⁴ Patients were randomized to receive one, none, or both treatments. Neither intervention showed significant change in total functional capacity (TFC), although patients treated with coenzyme Q10 showed a trend toward slowing in TFC decline (13%) over 30 months. In addition, there were beneficial trends on certain cognitive tests and tests of behavior in the coenzyme Q10-treated group. Interestingly, when both compounds were tested in two transgenic mouse models of HD, each compound was found to extend survival time and delay onset of motor deficits, cerebral atrophy, weight loss, and intranuclear inclusions.^{125,126} When the compounds were administered together they were found to have additive

beneficial effects. The discrepancy between mouse and human trial results highlights the difficulty in animal models of HD in predicting efficacy in humans. The CARE-HD trial was powered to detect a difference of 40% slowing; thus a 13% slowing is not large enough to be considered statistically significant.

Apoptosis/caspases

Although there is scant evidence for apoptosis in human HD, the demonstration that inhibiting caspase-1 could extend lifespan in transgenic HD mice gave credence to the idea that apoptosis was a legitimate target for therapeutics.⁵² Minocycline, a second-generation tetracycline, has been shown to inhibit caspase-1 and inducible NOS after experimental ischemia and to reduce infarct size.^{137,138} Minocycline was recently tested in transgenic HD mice and found to delay motoric decline (as measured by Rotarod performance) and extend survival time by 14%. Minocycline, however, did not affect weight, nor did it inhibit the formation of intranuclear inclusions or decrease in receptor binding. In addition, it was found that inhibition of both caspase-1 and caspase-3 was required for neuroprotection as drugs targeting one or the other did not extend survival.¹³⁹ As minocycline has been used in humans and known to have few side effects, it is a promising candidate for HD clinical trials. A phase II clinical trial assessing the safety and efficacy of minocycline in human HD patients has recently been completed. While minocycline is unlikely to be a cure for HD, in some ways it is the ideal first treatment; it is an inexpensive drug and has few side effects, and there are millions of patient-years of tolerability and toxicity experience. Minocycline may someday be one of the drugs that is used in a combinatorial "cocktail" approach, such as is currently used for HIV/Acquired Immune Deficiency Syndrome.

Cystamine, a caspase inhibitor, has also been studied in transgenic HD mice. Cystamine was originally tested as a therapeutic candidate given its ability to inhibit transglutaminases. Transglutaminases have been implicated as a possible mechanism of aggregate formation, by possibly cross-linking molecules of mutant huntingtin. Because mutant huntingtin aggregation appeared to correlate with disease phenotype, a transglutaminase inhibitor such as cystamine was tested for its potential ability to reduce aggregate formation and, in turn, extend survival. Both oral and intraperitoneal administration of cystamine extended survival in transgenic HD mice and reduced the number and size of aggregates^{140,141}. Recently, cystamine has been found to inhibit caspase-3 activity *in vitro*, suggesting that cystamine may work through a variety of mechanisms, including caspase inhibition, to prolong neuronal survival in HD.¹⁴²

Transcriptional dysregulation

The role of transcriptional dysregulation in HD has been an exciting area of research over the last few years.¹⁰⁸ Recent studies indicate that transcriptional repression is the main result of transcriptional dysregulation and could be explained by the recruitment and sequestering of transcription factors by mutant huntingtin. Because many of these nuclear factors are involved, directly or indirectly, in histone acetylation, neurotherapeutic research in this area has focused on histone deacetylase inhibitors (HDACs). The competing activities of histone acetyl transferases and histone deacetylases work to modify chromatin and thereby regulate transcription. Recent studies in cell culture, yeast, and *Drosophila* models of polyglutamine disease have shown that HDACs can reduce polyglutamine toxicity.^{143–145} A recent study tested the effects of suberoylanilide hydroxamic acid (SAHA) in a transgenic mouse model of HD.¹⁴⁶ Mice treated with SAHA demonstrated improved motor impairment and less striatal neuronal loss though there was no significant effect on weight or polyglutamine aggregation. The testing of other HDAC inhibitors in mouse models of HD is underway.

Restorative therapies

One obvious approach to alleviating the damage caused by dead striatal neurons is to transplant a new source of neurons. Various approaches, including porcine xenografts¹⁴⁷ and human fetal neurons^{148,149} have been tried. While there has not been definitive demonstration of clinical improvement with transplantation, a multi-center European trial is ongoing.¹⁴⁸ In one trial, an HD patient who had received a fetal cell transplant died of presumably unrelated causes 18 months later. Autopsy analysis demonstrated that the implanted cells had survived, expressed appropriate neurochemical markers, and received innervation by host dopaminergic cells, demonstrating that transplanted tissue could persist for 18 months.¹⁵⁰ As transplantation technology is improved, more advances will certainly be made. However, significant problems remain, not the least of which is that HD is a disease which affects multiple brain regions aside from the striatum.¹⁵¹

Various growth factors have been investigated in terms of their ability to maintain target cell populations. Activin-A, a nerve cell survival molecule, has also been tested in the quinolinic acid lesion model of HD.¹⁵² Quinolinic acid-treated rats that received intrastriatal infusion of activin-A displayed an improved phenotype with attenuation of cell death in a number of striatal nerve populations: striatal cholinergic interneurons and striatal projection neurons. Another approach has been to use genetically engineered cells that secrete various growth factors including ciliary neurotrophic factor, basic fibroblast growth factor, nerve growth factor, BDNF, NT3

and NT4.^{153–155} Interestingly, neurturin, a member of the GDNF growth factor family, demonstrated selective protection of enkephalin-containing striatal projections neurons, while GDNF selectively protected substance P-containing neurons in an excitotoxic lesion model of HD.¹⁵⁵

One potential drawback of the transplantation literature is that these experiments have largely relied on chemical lesion models of HD. To date, there has been no report of restoring growth factors to a transgenic mouse model of HD, although deficits in BDNF have been reported.⁵¹ There is one report of transplantation of anterior cingulate cortex into transgenic HD mice showing some motor improvement, pointing out the cortex as a possible target for transplantation.¹⁵⁶

Stem cells have garnered substantial interest as a potential source of restorative neurons.¹⁵⁷ A recent report points to an unsuspected source of neural stem cells: the Huntington's disease brain itself. Curtis et al.¹⁵⁸ found an increase in cell proliferation in the subependymal layer adjacent to the caudate nucleus of the human HD brain. Interestingly, these proliferative cells expressed both neuronal and glial cell markers. Furthermore, the degree of cell proliferation increased with pathological severity and with increasing CAG repeats in the *HD* gene. These observations raise the possibility that the HD brain itself could be coaxed into generating replacement neurons, especially if the rate of disease pathogenesis could be slowed by other means.

SUMMARY

Huntington's disease remains an important model disorder. The revolution in molecular genetics which led to discovery of the *HD* gene in 1993 has opened the door to discovery of basic pathogenic mechanisms. While the elucidation of multiple pathogenic mechanisms may be confusing, the other benefit is that such understanding permits development of multiple therapeutic targets. Looking forward, Huntington's disease is likely to remain an important model disease in neurotherapeutics.

Examination of human postmortem material as well as informative transgenic animal models has made it clear that although there is a single genetic cause, there are in fact numerous pathologic mechanisms that are unleashed in HD. While the multiplicity of disease mechanisms makes it unlikely that any single therapeutic approach will be successful, this complexity makes credible the idea that combination therapy could be successful. For example, while there have now been over 12 reports of slow disease progression in transgenic HD mice, recent studies indicate even more efficacy when two or more therapeutic agents are used in combination. "Rational therapy" therefore takes on a different connotation with respect to Huntington's disease. The first clinically suc-

cessful therapies will likely target downstream mechanisms such as glutamate excitotoxicity or apoptosis. In conjunction with this approach, more upstream mechanisms will also be tested, such as mitochondrial compromise or transcriptional dysregulation. In the future, innovative therapies such as selective disease allele inactivation or neural transplantation are also like to play a valuable role in attacking HD. A combinatorial cocktail approach targeting each of these important mechanisms is likely to yield, at long last, an effective therapy for this dreaded disease.

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